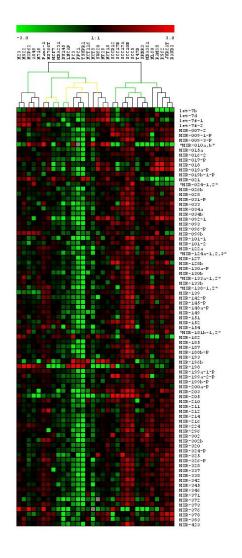
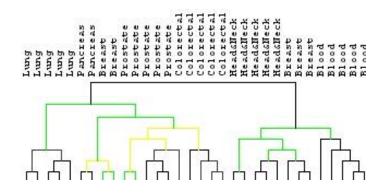
A B





Supplemental Figure 2. Heatmap of miRNA precursor expression in 35 human cancer cell lines, filtered data. The names of the cancer cell lines are listed on the top of the figure. The relative expression of each gene was determined as described in Materials and Methods. Unsupervised hierarchical clustering was performed on a subset of 92 genes which are differentially expressed (P = 0.01) among groups of cell lines as determined by ANOVA multi-group comparison test. The heatmap color legend is on the top. B., dendrogram of clustering data with the class of cell lines listed.